

<213> Artificial Sequence

<223> Description of Artificial Sequence:Fc IL-1

ANTAGONIST

<400> 1060

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Met Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu
 1              5              10              15

Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu
              20              25              30

Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser
              35              40              45

His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu
              50              55              60

Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr
 65              70              75              80

Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn
              85              90              95

Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro
              100              105              110

Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln
              115              120              125

Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val
 130              135              140

Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val
 145              150              155              160

Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro
              165              170              175

Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr
              180              185              190

Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val
              195              200              205

Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu
 210              215              220

Ser Pro Gly Lys Gly Gly Gly Gly Gly Phe Glu Trp Thr Pro Gly Tyr

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225

230

235

240

Trp Gln Pro Tyr Ala Leu Pro Leu

245

<210> 1061

<211> 757

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: IL-1 ANTAGONIST
Fc

<220>

<221> CDS

<222> (4) .. (747)

<400> 1061

cat atg ttc gaa tgg acc ccg ggt tac tgg cag ccg tac gct ctg ccg	48
Met Phe Glu Trp Thr Pro Gly Tyr Trp Gln Pro Tyr Ala Leu Pro	
1 5 10 15	
ctg ggt gga ggc ggt ggg gac aaa act cac aca tgt cca cct tgc cca	96
Leu Gly Gly Gly Gly Gly Asp Lys Thr His Thr Cys Pro Pro Cys Pro	
20 25 30	
gca cct gaa ctc ctg ggg gga ccg tca gtt ttc ctc ttc ccc cca aaa	144
Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys	
35 40 45	
ccc aag gac acc ctc atg atc tcc cgg acc cct gag gtc aca tgc gtg	192
Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val	
50 55 60	
gtg gtg gac gtg agc cac gaa gac cct gag gtc aag ttc aac tgg tac	240
Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr	
65 70 75	
gtg gac ggc gtg gag gtg cat aat gcc aag aca aag ccg cgg gag gag	288
Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu	
80 85 90 95	
cag tac aac agc acg tac cgt gtg gtc agc gtc ctc acc gtc ctg cac	336
Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His	
100 105 110	

cag gac tgg ctg aat ggc aag gag tac aag tgc aag gtc tcc aac aaa 384
 Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
 115 120 125

gcc ctc cca gcc ccc atc gag aaa acc atc tcc aaa gcc aaa ggg cag 432
 Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
 130 135 140

ccc cga gaa cca cag gtg tac acc ctg ccc cca tcc cgg gat gag ctg 480
 Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu
 145 150 155

acc aag aac cag gtc agc ctg acc tgc ctg gtc aaa ggc ttc tat ccc 528
 Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
 160 165 170 175

agc gac atc gcc gtg gag tgg gag agc aat ggg cag ccg gag aac aac 576
 Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
 180 185 190

tac aag acc acg cct ccc gtg ctg gac tcc gac ggc tcc ttc ttc ctc 624
 Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
 195 200 205

tac agc aag ctc acc gtg gac aag agc agg tgg cag cag ggg aac gtc 672
 Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
 210 215 220

ttc tca tgc tcc gtg atg cat gag gct ctg cac aac cac tac acg cag 720
 Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
 225 230 235

aag agc ctc tcc ctg tct ccg ggt aaa taatggatcc 757
 Lys Ser Leu Ser Leu Ser Pro Gly Lys
 240 245

<210> 1062

<211> 248

<212> PRT

<213> Artificial Sequence

 <223> Description of Artificial Sequence: IL-1 ANTAGONIST
 Fc

<400> 1062

 Met Phe Glu Trp Thr Pro Gly Tyr Trp Gln Pro Tyr Ala Leu Pro Leu
 1 5 10 15

Gly Gly Gly Gly Gly Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala
 20 25 30
 Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro
 35 40 45
 Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val
 50 55 60
 Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val
 65 70 75 80
 Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln
 85 90 95
 Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln
 100 105 110
 Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala
 115 120 125
 Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro
 130 135 140
 Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr
 145 150 155 160
 Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser
 165 170 175
 Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr
 180 185 190
 Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr
 195 200 205
 Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe
 210 215 220
 Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys
 225 230 235 240
 Ser Leu Ser Leu Ser Pro Gly Lys
 245

<210> 1063
 <211> 773
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Fc-VEGF
 ANTAGONIST

<220>

<221> CDS

<222> (4)..(759)

<400> 1063

cat atg gac aaa act cac aca tgt cca ccg tgc cca gca cct gaa ctc	48
Met Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu	
1 5 10 15	
ctg ggg gga ccg tca gtt ttc ctc ttc ccc cca aaa ccc aag gac acc	96
Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr	
20 25 30	
ctc atg atc tcc cgg acc cct gag gtc aca tgc gtg gtg gtg gac gtg	144
Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val	
35 40 45	
agc cac gaa gac cct gag gtc aag ttc aac tgg tac gtg gac ggc gtg	192
Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val	
50 55 60	
gag gtg cat aat gcc aag aca aag ccg cgg gag gag cag tac aac agc	240
Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser	
65 70 75	
acg tac cgt gtg gtc agc gtc ctc acc gtc ctg cac cag gac tgg ctg	288
Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu	
80 85 90 95	
aat ggc aag gag tac aag tgc aag gtc tcc aac aaa gcc ctc cca gcc	336
Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala	
100 105 110	
ccc atc gag aaa acc atc tcc aaa gcc aaa ggg cag ccc cga gaa cca	384
Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro	
115 120 125	
cag gtg tac acc ctg ccc cca tcc cgg gat gag ctg acc aag aac cag	432
Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln	

130	135	140	
gtc agc ctg acc tgc ctg gtc aaa ggc ttc tat ccc agc gac atc gcc			480
Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala			
145	150	155	
gtg gag tgg gag agc aat ggg cag ccg gag aac aac tac aag acc acg			528
Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr			
160	165	170	175
cct ccc gtg ctg gac tcc gac ggc tcc ttc ttc ctc tac agc aag ctc			576
Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu			
	180	185	190
acc gtg gac aag agc agg tgg cag cag ggg aac gtc ttc tca tgc tcc			624
Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser			
	195	200	205
gtg atg cat gag gct ctg cac aac cac tac acg cag aag agc ctc tcc			672
Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser			
	210	215	220
ctg tct ccg ggt aaa ggt ggt ggt ggt ggt gtt gaa ccg aac tgt gac			720
Leu Ser Pro Gly Lys Gly Gly Gly Gly Gly Val Glu Pro Asn Cys Asp			
	225	230	235
atc cat gtt atg tgg gaa tgg gaa tgt ttt gaa cgt ctg taactcgagg			769
Ile His Val Met Trp Glu Trp Glu Cys Phe Glu Arg Leu			
240	245	250	
atcc			773

<210> 1064

<211> 252

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence:Fc-VEGF

ANTAGONIST

<400> 1064

Met	Asp	Lys	Thr	His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu
1				5					10					15	

Gly	Gly	Pro	Ser	Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu
		20						25					30		

Met	Ile	Ser	Arg	Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

35	40	45
His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu		
50	55	60
Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr		
65	70	75
Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn		
	85	90
Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro		
	100	105
Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln		
	115	120
Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val		
	130	135
Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val		
145	150	155
Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro		
	165	170
Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr		
	180	185
Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val		
	195	200
Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu		
	210	215
Ser Pro Gly Lys Gly Gly Gly Gly Gly Val Glu Pro Asn Cys Asp Ile		
225	230	235
His Val Met Trp Glu Trp Glu Cys Phe Glu Arg Leu		
	245	250

<210> 1065

<211> 773

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: VEGF ANTAGONIST
Fc

<220>

<221> CDS

<222> (4) .. (759)

<400> 1065

cat atg gtt gaa ccg aac tgt gac atc cat gtt atg tgg gaa tgg gaa	48
Met Val Glu Pro Asn Cys Asp Ile His Val Met Trp Glu Trp Glu	
1 5 10 15	
tgt ttt gaa cgt ctg ggt ggt ggt ggt ggt gac aaa act cac aca tgt	96
Cys Phe Glu Arg Leu Gly Gly Gly Gly Gly Asp Lys Thr His Thr Cys	
20 25 30	
cca ccg tgc cca gca cct gaa ctc ctg ggg gga ccg tca gtt ttc ctc	144
Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu	
35 40 45	
ttc ccc cca aaa ccc aag gac acc ctc atg atc tcc cgg acc cct gag	192
Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu	
50 55 60	
gtc aca tgc gtg gtg gtg gac gtg agc cac gaa gac cct gag gtc aag	240
Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys	
65 70 75	
ttc aac tgg tac gtg gac ggc gtg gag gtg cat aat gcc aag aca aag	288
Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys	
80 85 90 95	
ccg cgg gag gag cag tac aac agc acg tac cgt gtg gtc agc gtc ctc	336
Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu	
100 105 110	
acc gtc ctg cac cag gac tgg ctg aat ggc aag gag tac aag tgc aag	384
Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys	
115 120 125	
gtc tcc aac aaa gcc ctc cca gcc ccc atc gag aaa acc atc tcc aaa	432
Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys	
130 135 140	
gcc aaa ggg cag ccc cga gaa cca cag gtg tac acc ctg ccc cca tcc	480
Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser	
145 150 155	

cgg gat gag ctg acc aag aac cag gtc agc ctg acc tgc ctg gtc aaa 528
 Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys
 160 165 170 175

 ggc ttc tat ccc agc gac atc gcc gtg gag tgg gag agc aat ggg cag 576
 Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln
 180 185 190

 ccg gag aac aac tac aag acc acg cct ccc gtg ctg gac tcc gac ggc 624
 Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly
 195 200 205

 tcc ttc ttc ctc tac agc aag ctc acc gtg gac aag agc agg tgg cag 672
 Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln
 210 215 220

 cag ggg aac gtc ttc tca tgc tcc gtg atg cat gag gct ctg cac aac 720
 Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn
 225 230 235

 cac tac acg cag aag agc ctc tcc ctg tct ccg ggt aaa taactcgagg 769
 His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
 240 245 250

 atcc 773

<210> 1066

<211> 252

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: VEGF ANTAGONIST

Fc

<400> 1066

Met Val Glu Pro Asn Cys Asp Ile His Val Met Trp Glu Trp Glu Cys
 1 5 10 15

Phe Glu Arg Leu Gly Gly Gly Gly Gly Asp Lys Thr His Thr Cys Pro
 20 25 30

Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe
 35 40 45

Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val
 50 55 60

Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe
 65 70 75 80
 Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro
 85 90 95
 Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr
 100 105 110
 Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val
 115 120 125
 Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala
 130 135 140
 Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg
 145 150 155 160
 Asp Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly
 165 170 175
 Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro
 180 185 190
 Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser
 195 200 205
 Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln
 210 215 220
 Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His
 225 230 235 240
 Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
 245 250

<210> 1067

<211> 748

<212> DNA

<213> Artificial Sequence

<220>

 <223> Description of Artificial Sequence:Fc-MMP
 INHIBITOR

<220>

<221> CDS

<222> (4) .. (732)

<400> 1067

cat atg gac aaa act cac aca tgt cca cct tgt cca gct ccg gaa ctc	48
Met Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu	
1 5 10 15	
ctg ggg gga ccg tca gtc ttc ctc ttc ccc cca aaa ccc aag gac acc	96
Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr	
20 25 30	
ctc atg atc tcc cgg acc cct gag gtc aca tgc gtg gtg gtg gac gtg	144
Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val	
35 40 45	
agc cac gaa gac cct gag gtc aag ttc aac tgg tac gtg gac ggc gtg	192
Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val	
50 55 60	
gag gtg cat aat gcc aag aca aag ccg cgg gag gag cag tac aac agc	240
Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser	
65 70 75	
acg tac cgt gtg gtc agc gtc ctc acc gtc ctg cac cag gac tgg ctg	288
Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu	
80 85 90 95	
aat ggc aag gag tac aag tgc aag gtc tcc aac aaa gcc ctc cca gcc	336
Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala	
100 105 110	
ccc atc gag aaa acc atc tcc aaa gcc aaa ggg cag ccc cga gaa cca	384
Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro	
115 120 125	
cag gtg tac acc ctg ccc cca tcc cgg gat gag ctg acc aag aac cag	432
Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln	
130 135 140	
gtc agc ctg acc tgc ctg gtc aaa ggc ttc tat ccc agc gac atc gcc	480
Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala	
145 150 155	
gtg gag tgg gag agc aat ggg cag ccg gag aac aac tac aag acc acg	528
Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr	
160 165 170 175	

cct ccc gtg ctg gac tcc gac ggc tcc ttc ttc ctc tac agc aag ctc 576
 Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu
 180 185 190

acc gtg gac aag agc agg tgg cag cag ggg aac gtc ttc tca tgc tcc 624
 Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser
 195 200 205

gtg atg cat gag gct ctg cac aac cac tac acg cag aag agc ctc tcc 672
 Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser
 210 215 220

ctg tct ccg ggt aaa ggt gga ggt ggt ggt tgc acc acc cac tgg ggt 720
 Leu Ser Pro Gly Lys Gly Gly Gly Gly Gly Cys Thr Thr His Trp Gly
 225 230 235

ttc acc ctg tgc taatggatcc ctcgag 748
 Phe Thr Leu Cys
 240

<210> 1068

<211> 243

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence:Fc-MMP
 INHIBITOR

<400> 1068

Met Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu
 1 5 10 15

Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu
 20 25 30

Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser
 35 40 45

His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu
 50 55 60

Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr
 65 70 75 80

Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn
 85 90 95

Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro

100	105	110
Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln 115	120	125
Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val 130	135	140
Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val 145	150	155 160
Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro 165	170	175
Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr 180	185	190
Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val 195	200	205
Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu 210	215	220
Ser Pro Gly Lys Gly Gly Gly Gly Gly Cys Thr Thr His Trp Gly Phe 225	230	235 240
Thr Leu Cys		

<210> 1069

<211> 763

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:MMP INHIBITOR
Fc

<220>

<221> CDS

<222> (4)..(753)

<400> 1069

cat atg tgc acc acc cac tgg ggt ttc acc ctg tgc ggt gga ggc ggt	48
Met Cys Thr Thr His Trp Gly Phe Thr Leu Cys Gly Gly Gly Gly	
1 5 10 15	

ggg gac aaa ggt gga ggc ggt ggg gac aaa act cac aca tgt cca cct	96
Gly Asp Lys Gly Gly Gly Gly Gly Asp Lys Thr His Thr Cys Pro Pro	
20 25 30	
tgc cca gca cct gaa ctc ctg ggg gga ccg tca gtt ttc ctc ttc ccc	144
Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro	
35 40 45	
cca aaa ccc aag gac acc ctc atg atc tcc cgg acc cct gag gtc aca	192
Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr	
50 55 60	
tgc gtg gtg gtg gac gtg agc cac gaa gac cct gag gtc aag ttc aac	240
Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn	
65 70 75	
tgg tac gtg gac ggc gtg gag gtg cat aat gcc aag aca aag ccg cgg	288
Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg	
80 85 90 95	
gag gag cag tac aac agc acg tac cgt gtg gtc agc gtc ctc acc gtc	336
Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val	
100 105 110	
ctg cac cag gac tgg ctg aat ggc aag gag tac aag tgc aag gtc tcc	384
Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser	
115 120 125	
aac aaa gcc ctc cca gcc ccc atc gag aaa acc atc tcc aaa gcc aaa	432
Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys	
130 135 140	
ggg cag ccc cga gaa cca cag gtg tac acc ctg ccc cca tcc cgg gat	480
Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp	
145 150 155	
gag ctg acc aag aac cag gtc agc ctg acc tgc ctg gtc aaa ggc ttc	528
Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe	
160 165 170 175	
tat ccc agc gac atc gcc gtg gag tgg gag agc aat ggg cag ccg gag	576
Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu	
180 185 190	
aac aac tac aag acc acg cct ccc gtg ctg gac tcc gac ggc tcc ttc	624
Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe	
195 200 205	

ttc ctc tac agc aag ctc acc gtg gac aag agc agg tgg cag cag ggg 672
 Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly
 210 215 220

aac gtc ttc tca tgc tcc gtg atg cat gag gct ctg cac aac cac tac 720
 Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr
 225 230 235

acg cag aag agc ctc tcc ctg tct ccg ggt aaa taatggatcc 763
 Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
 240 245 250

<210> 1070

<211> 250

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence:MMP INHIBITOR
 Fc

<400> 1070

Met Cys Thr Thr His Trp Gly Phe Thr Leu Cys Gly Gly Gly Gly Gly
 1 5 10 15

Asp Lys Gly Gly Gly Gly Gly Asp Lys Thr His Thr Cys Pro Pro Cys
 20 25 30

Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro
 35 40 45

Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys
 50 55 60

Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp
 65 70 75 80

Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu
 85 90 95

Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu
 100 105 110

His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn
 115 120 125

Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly
 130 135 140

Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu
145 150 155 160

Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr
165 170 175

Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn
180 185 190

Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe
195 200 205

Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn
210 215 220

Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr
225 230 235 240

Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
245 250

<210> 1071

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:INTEGRIN
BINDING PEPTIDE

<400> 1071

Cys Gly Arg Glu Cys Pro Arg Leu Cys Gln Ser Ser Cys
1 5 10

<210> 1072

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:INTEGRIN
BINDING PEPTIDE

<400> 1072

Cys Asn Gly Arg Cys Val Ser Gly Cys Ala Gly Arg Cys
1 5 10

<210> 1073

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:INTEGRIN
BINDING PEPTIDE

<400> 1073

Cys Leu Ser Gly Ser Leu Ser Cys
1 5

<210> 1074

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:INTEGRIN
BINDING PEPTIDE

<400> 1074

Asn Gly Arg Ala His Ala
1 5

<210> 1075

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:INTEGRIN
BINDING PEPTIDE

<220>

<221> CDS

<222> (10)..(189)

<400> 1075

Cys Asn Gly Arg Cys

1

5

<210> 1076

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:INTEGRIN
BINDING PEPTIDE

<400> 1076

Cys Asp Cys Arg Gly Asp Cys Phe Cys

1

5

<210> 1077

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:INTEGRIN
BINDING PEPTIDE

<400> 1077

Cys Gly Ser Leu Val Arg Cys

1

5

<210> 1078

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:INTEGRIN
BINDING PEPTIDE

<400> 1078

Arg Thr Asp Leu Asp Ser Leu Arg

1

5

<210> 1079

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:INTEGRIN
BINDING PEPTIDE

<400> 1079 .

Gly Asp Leu Asp Leu Leu Lys Leu Arg Leu Thr Leu

1

5

10

<210> 1080

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial
Sequence:INTEGRIN-BINDING PEPTIDE

<400> 1080

Gly Asp Leu His Ser Leu Arg Gln Leu Leu Ser Arg

1

5

10

<210> 1081

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial
Sequence:INTEGRIN-BINDING PEPTIDE

<400> 1081

Arg Asp Asp Leu His Met Leu Arg Leu Gln Leu Trp

1

5

10

<210> 1082

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial

Sequence:INTEGRIN-BINDING PEPTIDE

<400> 1082

Ser Ser Asp Leu His Ala Leu Lys Lys Arg Tyr Gly

1

5

10

<210> 1083

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial

Sequence:INTEGRIN-BINDING PEPTIDE

<400> 1083

Arg Gly Asp Leu Lys Gln Leu Ser Glu Leu Thr Trp

1

5

10

<210> 1084

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial

Sequence:INTEGRIN-BINDING PEPTIDE

<400> 1084

Arg Gly Asp Leu Ala Ala Leu Ser Ala Pro Pro Val

1

5

10

<210> 1085

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:TNF-ANTAGONIST
PEPTIDE

<400> 1085

Asp	Phe	Leu	Pro	His	Tyr	Lys	Asn	Thr	Ser	Leu	Gly	His	Arg	Pro
1				5				10					15	

<210> 1086

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:VEGF ANTAGONIST
PEPTIDE

<400> 1086

Gly	Glu	Arg	Trp	Cys	Phe	Asp	Gly	Pro	Leu	Thr	Trp	Val	Cys	Gly	Glu
1				5				10					15		

Glu Ser

<210> 1087

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:VEGF ANTAGONIST
PEPTIDE

<400> 1087

Arg	Gly	Trp	Val	Glu	Ile	Cys	Val	Ala	Asp	Asp	Asn	Gly	Met	Cys	Val
1				5				10				15			

Thr Glu Ala Gln

... 20

<210> 1088
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:VEGF ANTAGONIST
PEPTIDE

<400> 1088
Gly Trp Asp Glu Cys Asp Val Ala Arg Met Trp Glu Trp Glu Cys Phe
1 5 10 15

Ala Gly Val

<210> 1089
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:VEGF ANTAGONIST
PEPTIDE

<400> 1089
Arg Gly Trp Val Glu Ile Cys Glu Ser Asp Val Trp Gly Arg Cys Leu
1 5 10 15

<210> 1090
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:VEGF ANTAGONIST
PEPTIDE

<400> 1090
Arg Gly Trp Val Glu Ile Cys Glu Ser Asp Val Trp Gly Arg Cys Leu
1 5 10 15

<210> 1091
<211> 19
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:VEGF ANTAGONIST
PEPTIDE

<400> 1091

Gly	Gly	Asn	Glu	Cys	Asp	Ile	Ala	Arg	Met	Trp	Glu	Trp	Glu	Cys	Phe
1				5				10						15	

Glu Arg Leu

<210> 1092
<211> 16
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:VEGF ANTAGONIST
PEPTIDE

<400> 1092

Arg	Gly	Trp	Val	Glu	Ile	Cys	Ala	Ala	Asp	Asp	Tyr	Gly	Arg	Cys	Leu
1				5					10					15	

<210> 1093
<211> 8
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:MMP INHIBITOR
PEPTIDE

<400> 1093

Cys	Leu	Arg	Ser	Gly	Xaa	Gly	Cys
1				5			

<210> 1094
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:MMP INHIBITOR
PEPTIDE

<400> 1094
Cys Xaa Xaa His Trp Gly Phe Xaa Xaa Cys
1 5 10

<210> 1095
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:MMP INHIBITOR
PEPTIDE

<400> 1095
Cys Xaa Pro Xaa Cys
1 5

<210> 1096
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:MMP INHIBITOR
PEPTIDE

<400> 1096
Cys Arg Arg His Trp Gly Phe Glu Phe Cys
1 5 10

<210> 1097
<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:MMP INHIBITOR
PEPTIDE

<400> 1097

Ser	Thr	Thr	His	Trp	Gly	Phe	Thr	Leu	Ser
1				5					10

<210> 1098

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:MMP INHIBITOR
PEPTIDE

<400> 1098

Cys	Ser	Leu	His	Trp	Gly	Phe	Trp	Trp	Cys
1				5					10

<210> 1099

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:CARBOHYDRATE
(GD1 ALPHA) MIMETIC PEPTIDE

<400> 1099

Trp	His	Trp	Arg	His	Arg	Ile	Pro	Leu	Gln	Leu	Ala	Ala	Gly	Arg
1				5					10					15

<210> 1100

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BETA-2 GP1AB
BINDING PEPTIDE

<400> 1100

Leu Lys Thr Pro Arg Val
1 5

<210> 1101

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BETA-2 GP1AB
BINDING PEPTIDE

<400> 1101

Asn Thr Leu Lys Thr Pro Arg Val
1 5

<210> 1102

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BETA-2 GP1AB
BINDING PROTEIN

<400> 1102

Asn Thr Leu Lys Thr Pro Arg Val Gly Gly Cys
1 5 10

<210> 1103

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BETA-2 GP1AB
BINDING PROTEIN

<400> 1103

Lys Asp Lys Ala Thr Phe
1 5

<210> 1104

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BETA-1 GP1AB
BINDING PROTEIN

<400> 1104

Lys Asp Lys Ala Thr Phe Gly Cys His Asp
1 5 10

<210> 1105

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BETA-2 GP1AB
BINDING PEPTIDE

<400> 1105

Lys Asp Lys Ala Thr Phe Gly Cys His Asp Gly Cys
1 5 10

<210> 1106

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BETA-2 GP1AB
BINDING PROTEIN

<400> 1106

Thr Leu Arg Val Tyr Lys

1

5

<210> 1107

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BETA-2 GP1AB
BINDING PROTEIN

<400> 1107

Ala Thr Leu Arg Val Tyr Lys Gly Gly

1

5

<210> 1108

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BETA-2 GP1AB
BINDING PROTEIN

<400> 1108

Cys Ala Thr Leu Arg Val Tyr Lys Gly Gly

1

5

10

<210> 1109

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: MEMBRANE
TRANSPORTING PEPTIDE

<400> 1109

Ile Asn Leu Lys Ala Leu Ala Ala Leu Ala Lys Lys Ile Leu

1

5

10

<210> 1110
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:MEMBRANE
TRANSPORTING PEPTIDE

<400> 1110
Gly Trp Thr Leu Asn Ser Ala Gly Tyr Leu Leu Gly
1 5 10

<210> 1111
<211> 27
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:MEMBRANE
TRANSPORTING PEPTIDE

<400> 1111
Gly Trp Thr Leu Asn Ser Ala Gly Tyr Leu Leu Gly Lys Ile Asn Leu
1 5 10 15
Lys Ala Leu Ala Ala Leu Ala Lys Lys Ile Leu
20 25

<210> 1112
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:FC PCR PRIMER

<400> 1112
aacataagta cctgtaggat cg

22

<210> 1113
<211> 81

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Fc-TNF ALPHA
PCR PRIMER

<220>

<221> CDS

<222> (1)..(126)

<400> 1113

ccg	cgg	atc	cat	tac	gga	cgg	tga	ccc	aga	gag	gtg	ttt	ttg	tag	tgc	48
Pro	Arg	Ile	His	Tyr	Gly	Arg		Pro	Arg	Glu	Val	Phe	Leu		Cys	
1				5				10					15			

ggc	agg	aag	tca	cca	cca	cct	cca	cct	tta	ccc	81
Gly	Arg	Lys	Ser	Pro	Pro	Pro	Pro	Pro	Leu	Pro	
			20				25				

<210> 1114

<211> 7

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence:Fc-TNF ALPHA
PCR PRIMER

<400> 1114

Pro	Arg	Ile	His	Tyr	Gly	Arg
1				5		

<210> 1115

<211> 6

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence:Fc-TNF ALPHA
PCR PRIMER

<400> 1115

Pro	Arg	Glu	Val	Phe	Leu
1				5	

<210> 1116

<211> 12

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence:Fc-TNF ALPHA
PCR PRIMER

<400> 1116

Cys Gly Arg Lys Ser Pro Pro Pro Pro Pro Leu Pro
1 5 10

<210> 1117

<211> 81

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:TNF-ALPHA
INHIBITOR-Fc PCR PRIMER

<400> 1117

gaataacata tggacttcct gccgcactac aaaaacacct ctctgggtca ccgtccgggt 60
ggaggcggtg gggacaaaac t 81

<210> 1118

<211> 81

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:IL-1 ANTAGONIST
PCR PRIMER

<400> 1118

ccgcggatcc attacagcgg cagagcgtac ggctgccagt aaccgggggt ccattcgaaa 60
ccaccacctc cacctttacc c 81

<210> 1119

<211> 81

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:IL-1 ANTAGONIST
-Fc PCR PRIMER

<400> 1119

gaataacata tggttcgaatg gacccccgggt tactggcagc cgtacgctct gccgctgggt 60
ggaggcgggtg gggacaaaac t 81

<210> 1120
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Fc-VEGF
ANTAGONIST OLIGONUCLEOTIDE

<400> 1120
gttgaaccga actgtgacat ccatgttatg tgggaatggg aatgttttga acgtctg 57

<210> 1121
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Fc-VEGF
ANTAGONIST OLIGONUCLEOTIDE

<400> 1121
cagacgttca aaacattccc attcccacat aacatggatg tcacagttcg gttcaac 57

<210> 1122
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Fc-VEGF
ANTAGONIST PCR TEMPLATE

<400> 1122
gttgaaccga actgtgacat ccatgttatg tgggaatggg aatgttttga acgtctg 57

<210> 1123
<211> 48
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Fc PRIMER

<400> 1123

atttgattct agaaggagga ataacatatg gacaaaactc acacatgt

48

<210> 1124

<211> 51

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Fc PRIMER

<400> 1124

gtcacagttc ggttcaacac caccaccacc acctttaccc ggagacaggg a

51

<210> 1125

<211> 54

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Fc-VEGF
ANTAGONIST PCR PRIMER

<400> 1125

tccctgtctc cgggtaaagg tggtggtggt ggtggtgaac cgaactgtga catc

54

<210> 1126

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Fc-VEGF
ANTAGONIST-Fc PCR PRIMER

<400> 1126

ccgcggatcc tcgagttaca gacgttcaaa acattccca

39

<210> 1127--

<211> 48

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Fc-VEGF
ANTAGONIST-Fc PCR PRIMER

<400> 1127

atttgattct agaaggagga ataacatatg gttgaaccga actgtgac

48

<210> 1128

<211> 51

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Fc-VEGF
ANTAGONIST-Fc PCR PRIMER

<400> 1128

acatgtgtga gttttgtcac caccaccacc acccagacgt tcaaaacatt c

51

<210> 1129

<211> 51

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Fc-PCR PRIMER

<400> 1129

gaatgttttg aacgtctggg tgggtggtggt ggtgacaaaa ctcacacatg t

51

<210> 1130

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Fc PCR PRIMER

<400> 1130

ccgcggatcc tcgagttatt taccgggaga caggagag

39

<210> 1131

<211> 66
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:Fc-MMP
 INHIBITOR PCR PRIMER

<400> 1131
 ccgcggatcc attagcacag ggtgaaaccc cagtgggtgg tgcaaccacc acctccacct 60
 ttaccc 66

<210> 1132
 <211> 63
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:MMP
 INHIBITOR-Fc PCR PRIMER

<400> 1132
 gaataacata tgtgcaccac ccactggggt ttcaccctgt gcggtggagg cggtggggac. 60
 aaa 63

<210> 1133
 <211> 20
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:Fc-TMP

<400> 1133
 Lys Gly Gly Gly Gly Gly Ile Glu Gly Pro Thr Leu Arg Gln Trp Leu
 1 5 10 15

Ala Ala Arg Ala
 20